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Sequence Listing was accepted with existing errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Wed Jun 20 10:43:59 EDT 2007
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Validated By CRFValidator v 1.0.2

Application No: 10715868   Version No: 2.1

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Finished: 2007-06-20 10:43:18.236
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Total Errors: 0
No. of SeqIDs Defined: 9
Actual SeqID Count: 9

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<110>   Arroyo, Juan
        Miller , Charles
        Catalan, John Avram
        Monath, Thomas P.

<120>   West Nile Virus Vaccine

<130>   06132/075002

<140>   US 10/715,868
<141>   2003-11-17

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                                        Met Ser Lys Lys Pro Gly
                                        1               5


ggg ccc ggc aag agc cgg gct gtc aat atg cta aaa cgc gga atg ccc      162
Gly Pro Gly Lys Ser Arg Ala Val Asn Met Leu Lys Arg Gly Met Pro
            10                  15                  20


cgc gtg ttg tcc ttg att gga ctg aag agg gct atg ttg agc ctg atc      210
Arg Val Leu Ser Leu Ile Gly Leu Lys Arg Ala Met Leu Ser Leu Ile
        25                  30                  35


gac ggc aag ggg cca ata cga ttt gtg ttg gct ctc ttg gcg ttc ttc      258
Asp Gly Lys Gly Pro Ile Arg Phe Val Leu Ala Leu Leu Ala Phe Phe
    40                  45                  50


agg ttc aca gca att gct ccg acc cga gca gtg ctg gat cga tgg aga      306
Arg Phe Thr Ala Ile Ala Pro Thr Arg Ala Val Leu Asp Arg Trp Arg
55                  60                  65                  70


ggt gtg aac aaa caa aca gcg atg aaa cac ctt ctg agt ttt aag aag      354
Gly Val Asn Lys Gln Thr Ala Met Lys His Leu Leu Ser Phe Lys Lys
                75                  80                  85


gaa cta ggg acc ttg acc agt gct atc aat cgg cgg agc tca aaa caa      402
Glu Leu Gly Thr Leu Thr Ser Ala Ile Asn Arg Arg Ser Ser Lys Gln
            90                  95                  100


aag aaa aga gga gga aag acc gga att gca gtc atg att ggc ctg atc      450
Lys Lys Arg Gly Gly Lys Thr Gly Ile Ala Val Met Ile Gly Leu Ile
        105                 110                 115


gcc agc gta gga gca gtt acc ctc tct aac ttc caa ggg aag gtg atg      498
Ala Ser Val Gly Ala Val Thr Leu Ser Asn Phe Gln Gly Lys Val Met
    120                 125                 130


atg acg gta aat gct act gac gtc aca gat gtc atc acg att cca aca      546
Met Thr Val Asn Ala Thr Asp Val Thr Asp Val Ile Thr Ile Pro Thr
135                 140                 145                 150


gct gct gga aag aac cta tgc att gtc aga gca atg gat gtg gga tac      594
Ala Ala Gly Lys Asn Leu Cys Ile Val Arg Ala Met Asp Val Gly Tyr
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atg tgc gat gat act atc act tat gaa tgc cca gtg ctg tcg gct ggt      642
Met Cys Asp Asp Thr Ile Thr Tyr Glu Cys Pro Val Leu Ser Ala Gly
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aat gat cca gaa gac atc gac tgt tgg tgc aca aag tca gca gtc tac      690
Asn Asp Pro Glu Asp Ile Asp Cys Trp Cys Thr Lys Ser Ala Val Tyr
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Arg Ser Leu Thr Val Gln Thr His Gly Glu Ser Thr Leu Ala Asn Lys
215              220              225              230


aag ggg gct tgg atg gac agc acc aag gcc aca agg tat ttg gta aaa        834
Lys Gly Ala Trp Met Asp Ser Thr Lys Ala Thr Arg Tyr Leu Val Lys
                 235              240              245


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Thr Glu Ser Trp Ile Leu Arg Asn Pro Gly Tyr Ala Leu Val Ala Ala
                 250              255              260


gtc att ggt tgg atg ctt ggg agc aac acc atg cag aga gtt gtg ttt        930
Val Ile Gly Trp Met Leu Gly Ser Asn Thr Met Gln Arg Val Val Phe
         265              270              275


gtc gtg cta ttg ctt ttg gtg gcc cca gct tac agc ttc aac tgc ctt        978
Val Val Leu Leu Leu Leu Val Ala Pro Ala Tyr Ser Phe Asn Cys Leu
         280              285              290


gga atg agc aac aga gac ttc ttg gaa gga gtg tct gga gca aca tgg       1026
Gly Met Ser Asn Arg Asp Phe Leu Glu Gly Val Ser Gly Ala Thr Trp
295              300              305              310


gtg gat ttg gtt ctc gaa ggc gac agc tgc gtg act atc atg tct aag       1074
Val Asp Leu Val Leu Glu Gly Asp Ser Cys Val Thr Ile Met Ser Lys
                 315              320              325


gac aag cct acc atc gat gtg aag atg atg aat atg gag gcg gcc aac       1122
Asp Lys Pro Thr Ile Asp Val Lys Met Met Asn Met Glu Ala Ala Asn
                 330              335              340


ctg gca gag gtc cgc agt tat tgc tat ttg gct acc gtc agc gat ctc       1170
Leu Ala Glu Val Arg Ser Tyr Cys Tyr Leu Ala Thr Val Ser Asp Leu
         345              350              355


tcc acc aaa gct gcg tgc ccg acc atg gga gaa gct cac aat gac aaa       1218
Ser Thr Lys Ala Ala Cys Pro Thr Met Gly Glu Ala His Asn Asp Lys
         360              365              370


cgt gct gac cca gct ttt gtg tgc aga caa gga gtg gtg gac agg ggc       1266
Arg Ala Asp Pro Ala Phe Val Cys Arg Gln Gly Val Val Asp Arg Gly
375              380              385              390


tgg ggc aac ggc tgc gga cta ttt ggc aaa gga agc att gac aca tgc       1314
Trp Gly Asn Gly Cys Gly Leu Phe Gly Lys Gly Ser Ile Asp Thr Cys
                 395              400              405


gcc aaa ttt gcc tgc tct acc aag gca ata gga aga acc atc ttg aaa       1362
Ala Lys Phe Ala Cys Ser Thr Lys Ala Ile Gly Arg Thr Ile Leu Lys
                 410              415              420


gag aat atc aag tac gaa gtg gcc att ttt gtc cat gga cca act act       1410
Glu Asn Ile Lys Tyr Glu Val Ala Ile Phe Val His Gly Pro Thr Thr
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gtg gag tcg cac gga aac tac tcc aca cag gtt gga gcc act cag gca      1458
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ggg aga ttc agc atc act cct gcg gcg cct tca tac aca cta aag ctt      1506
Gly Arg Phe Ser Ile Thr Pro Ala Ala Pro Ser Tyr Thr Leu Lys Leu
455                     460                     465                470

gga gaa tat gga gag gtg aca gtg gac tgt gaa cca cgg tca ggg att      1554
Gly Glu Tyr Gly Glu Val Thr Val Asp Cys Glu Pro Arg Ser Gly Ile
                   475                     480                     485

gac acc aat gca tac tac gtg atg act gtt gga aca aag acg ttc ttg      1602
Asp Thr Asn Ala Tyr Tyr Val Met Thr Val Gly Thr Lys Thr Phe Leu
                   490                     495                     500

gtc cat cgt gag tgg ttc atg gac ctc aac ctc cct tgg agc agt gct      1650
Val His Arg Glu Trp Phe Met Asp Leu Asn Leu Pro Trp Ser Ser Ala
         505                     510                     515

gga agt act gtg tgg agg aac aga gag acg tta atg gag ttt gag gaa      1698
Gly Ser Thr Val Trp Arg Asn Arg Glu Thr Leu Met Glu Phe Glu Glu
         520                     525                     530

cca cac gcc acg aag cag tct gtg ata gca ttg ggc tca caa gag gga      1746
Pro His Ala Thr Lys Gln Ser Val Ile Ala Leu Gly Ser Gln Glu Gly
535                     540                     545                550

gct ctg cat caa gct ttg gct gga gcc att cct gtg gaa ttt tca agc      1794
Ala Leu His Gln Ala Leu Ala Gly Ala Ile Pro Val Glu Phe Ser Ser
                   555                     560                     565

aac act gtc aag ttg acg tcg ggt cat ttg aag tgt aga gtg aag atg      1842
Asn Thr Val Lys Leu Thr Ser Gly His Leu Lys Cys Arg Val Lys Met
                   570                     575                     580

gaa aaa ttg cag ttg aag gga aca acc tat ggc gtc tgt tca aag gct      1890
Glu Lys Leu Gln Leu Lys Gly Thr Thr Tyr Gly Val Cys Ser Lys Ala
         585                     590                     595

ttc aag ttt ctt ggg act ccc gca gac aca ggt cac ggc act gtg gtg      1938
Phe Lys Phe Leu Gly Thr Pro Ala Asp Thr Gly His Gly Thr Val Val
         600                     605                     610

ttg gaa ttg cag tac act ggc acg gat gga cct tgc aaa gtt cct atc      1986
Leu Glu Leu Gln Tyr Thr Gly Thr Asp Gly Pro Cys Lys Val Pro Ile
615                     620                     625                630

tcg tca gtg gct tca ttg aac gac cta acg cca gtg ggc aga ttg gtc      2034
Ser Ser Val Ala Ser Leu Asn Asp Leu Thr Pro Val Gly Arg Leu Val
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act gtc aac cct ttt gtt tca gtg gcc acg gcc aac gct aag gtc ctg      2082
Thr Val Asn Pro Phe Val Ser Val Ala Thr Ala Asn Ala Lys Val Leu
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Ile Glu Leu Glu Pro Pro Phe Gly Asp Ser Tyr Ile Val Val Gly Arg
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gga gaa caa cag atc aat cac cat tgg cac aag tct gga agc agc att          2178
Gly Glu Gln Gln Ile Asn His His Trp His Lys Ser Gly Ser Ser Ile
        680                 685                 690

ggc aaa gcc ttt aca acc acc ctc aaa gga gcg cag aga cta gcc gct          2226
Gly Lys Ala Phe Thr Thr Thr Leu Lys Gly Ala Gln Arg Leu Ala Ala
695                 700                 705                 710

cta gga gac aca gct tgg gac ttt gga tca gtt gga ggg gtg ttc acc          2274
Leu Gly Asp Thr Ala Trp Asp Phe Gly Ser Val Gly Gly Val Phe Thr
                715                 720                 725

tca gtt ggg aag gct gtc cat caa gtg ttc gga gga gca ttc cgc tca          2322
Ser Val Gly Lys Ala Val His Gln Val Phe Gly Gly Ala Phe Arg Ser
                730                 735                 740

ctg ttc gga ggc atg tcc tgg ata acg caa gga ttg ctg ggg gct ctc          2370
Leu Phe Gly Gly Met Ser Trp Ile Thr Gln Gly Leu Leu Gly Ala Leu
                745                 750                 755

ctg ttg tgg atg ggc atc aat gct cgt gat agg tcc ata gct ctc acg          2418
Leu Leu Trp Met Gly Ile Asn Ala Arg Asp Arg Ser Ile Ala Leu Thr
        760                 765                 770

ttt ctc gca gtt gga gga gtt ctg ctc ttc ctc tcc gtg aac gtg cac          2466
Phe Leu Ala Val Gly Gly Val Leu Leu Phe Leu Ser Val Asn Val His
775                 780                 785                 790

gct gac act ggg tgt gcc ata gac atc agc cgg caa gag ctg aga tgt          2514
Ala Asp Thr Gly Cys Ala Ile Asp Ile Ser Arg Gln Glu Leu Arg Cys
                795                 800                 805

gga agt gga gtg ttc ata cac aat gat gtg gag gct tgg atg gac cgg          2562
Gly Ser Gly Val Phe Ile His Asn Asp Val Glu Ala Trp Met Asp Arg
                810                 815                 820

tac aag tat tac cct gaa acg cca caa ggc cta gcc aag atc att cag          2610
Tyr Lys Tyr Tyr Pro Glu Thr Pro Gln Gly Leu Ala Lys Ile Ile Gln
                825                 830                 835

aaa gct cat aag gaa gga gtg tgc ggt cta cga tca gtt tcc aga ctg          2658
Lys Ala His Lys Glu Gly Val Cys Gly Leu Arg Ser Val Ser Arg Leu
        840                 845                 850

gag cat caa atg tgg gaa gca gtg aag gac gag ctg aac act ctt ttg          2706
Glu His Gln Met Trp Glu Ala Val Lys Asp Glu Leu Asn Thr Leu Leu
855                 860                 865                 870

aag gag aat ggt gtg gac ctt agt gtc gtg gtt gag aaa cag gag gga          2754
Lys Glu Asn Gly Val Asp Leu Ser Val Val Val Glu Lys Gln Glu Gly
                875                 880                 885
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atg tac aag tca gca cct aaa cgc ctc acc gcc acc acg gaa aaa ttg        2802
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gaa att ggc tgg aag gcc tgg gga aag agt att tta ttt gca cca gaa        2850
Glu Ile Gly Trp Lys Ala Trp Gly Lys Ser Ile Leu Phe Ala Pro Glu
        905                 910                 915

ctc gcc aac aac acc ttt gtg gtt gat ggt ccg gag acc aag gaa tgt        2898
Leu Ala Asn Asn Thr Phe Val Val Asp Gly Pro Glu Thr Lys Glu Cys
        920                 925                 930

ccg act cag aat cgc gct tgg aat agc tta gaa gtg gag gat ttt gga        2946
Pro Thr Gln Asn Arg Ala Trp Asn Ser Leu Glu Val Glu Asp Phe Gly
935                 940                 945                 950

ttt ggt ctc acc agc act cgg atg ttc ctg aag gtc aga gag agc aac        2994
Phe Gly Leu Thr Ser Thr Arg Met Phe Leu Lys Val Arg Glu Ser Asn
                955                 960                 965

aca act gaa tgt gac tcg aag atc att gga acg gct gtc aag aac aac        3042
Thr Thr Glu Cys Asp Ser Lys Ile Ile Gly Thr Ala Val Lys Asn Asn
                970                 975                 980

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Leu Ala Ile His Ser Asp Leu Ser Tyr Trp Ile Glu Ser Arg Leu Asn
        985                 990                 995

gat acg  tgg aag ctt gaa agg  gca gtt ctg ggt gaa  gtc aaa tca        3135
Asp Thr  Trp Lys Leu Glu Arg  Ala Val Leu Gly Glu  Val Lys Ser
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Cys Thr  Trp Pro Glu Thr His  Thr Leu Trp Gly Asp  Gly Ile Leu
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gag agt  gac ttg ata ata cca  gtc aca ctg gcg gga  cca cga agc        3225
Glu Ser  Asp Leu Ile Ile Pro  Val Thr Leu Ala Gly  Pro Arg Ser
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aat cac  aat cgg aga cct ggg  tac aag aca caa aac  cag ggc cca        3270
Asn His  Asn Arg Arg Pro Gly  Tyr Lys Thr Gln Asn  Gln Gly Pro
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Trp Asp  Glu Gly Arg Val Glu  Ile Asp Phe Asp Tyr  Cys Pro Gly
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act acg  gtc acc ctg agt gag  agc tgc gga cac cgt  gga cct gcc        3360
Thr Thr  Val Thr Leu Ser Glu  Ser Cys Gly His Arg  Gly Pro Ala
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Thr Arg  Thr Thr Thr Glu Ser  Gly Lys Leu Ile Thr  Asp Trp Cys
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Gly Cys  Trp
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